

AMENDMENTS TO THE SPECIFICATION

Replace the paragraph beginning at page 87, line 11, which starts with "To further examine" with the following **amended** paragraph:

To further examine the relevance of the RelB:p52 selective binding site, we conducted a pattern search with two strings, namely AGGAGATTTG (*Elc*-κB) (SEQ ID NO:60) and GGGAGATTTG (*Blc*-κB) (SEQ ID NO:59) using the Trafac server and the BlastZ algorithm ([Http://bio.cse.psu.edu](http://bio.cse.psu.edu)). Closely similar (at least 8/10 identity) sites were detected within 5 kb upstream to the start sites of the *Sdf-1* and *Baff* genes, whose expression is known to be *Ikkα*-dependent (Dejardin *et al.* (2002) *Immunity*, 17, 525-535) (Figs 26D, 27A, 30D). We also detected similar and evolutionary conserved sites with the same region of several other genes, whose IKKα-dependence was previously unknown (Fig 30D). RT-PCR analysis revealed that two of these genes, *Rxra* and *Irf3*, coding for important transcription factors, were induced in stromal cells in response to anti-LTβR in a manner dependent on IKKα (Fig 30E).

Insert the paper copy of the Sequence Listing (enclosed) immediately prior to the claims that begin on page 89 of the originally filed specification.

Renumber the Claims pages consecutively after the Sequence Listing.

Renumber the Abstract page.